PCT

RAW SEQUENCE LISTING DATE: 07/08/2004
PATENT APPLICATION: **US/10/500,586** TIME: 16:27:07

Input Set : **A:\280277 Sequence Listings.txt**
Output Set: **N:\CRF4\07082004\J500586.raw**

<110> APPLICANT: KIM, Bum-Joon
BIOMEDLAB CORPORATION
<120> TITLE OF INVENTION: PRIMERS FOR AMPLIFYING HSP 65 GENE OF MYCOBACTERIAL SPECIES, HSP
65 GENE FRAGMENTS AND METHOD OF IDENTIFYING MYCOBACTERIAL SPECIES
WITH THE SAME
<130> FILE REFERENCE: OPP021096KR
C--> 10 **<140> CURRENT APPLICATION NUMBER: US/10/500,586**
C--> 10 **<141> CURRENT FILING DATE: 2004-06-30**
<150> PRIOR APPLICATION NUMBER: KR 10-2002-0004297
<151> PRIOR FILING DATE: 2002-01-24
<150> PRIOR APPLICATION NUMBER: KR 10-2002-0011648
<151> PRIOR FILING DATE: 2002-03-05
<160> NUMBER OF SEQ ID NOS: 56
<170> SOFTWARE: KopatentIn 1.71
<210> SEQ ID NO: 1
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<213> ORGANISM: Mycobacterium abscessus
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<210> SEQ ID NO: 2
<211> LENGTH: 604
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<213> ORGANISM: Mycobacterium africanum
<400> SEQUENCE: 2

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90 ggccgtcgag aaggtcaccc agaccctgct cagctcggcc aaggacgtcg agaccaagga 240
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117 aggtctgcgc aacgtcgctg ccggcgccaa cccgctcggc ctgaagcgcg gcatcgagaa 180
119 ggccgtcgag aagatcaccg agacgctcct caagagcgcc aaggaggtcg agaccaagga 240
121 ccagatcgcg gccaccgccg ggatctccgc gggcgaccag accatcggtg acctgatcgc 300
123 cgaggccatg gacaaggtcg gcaacgaggg tgtcatcacc gtcgaggagt cgaacacctt 360
125 cggcctgcag ctcgagctca ccgagggtat gcgcttcgac aagggctaca tctcgggtta 420
127 cttcgtgacc gacgccgagc gtcaggaagc ggtcctcgag gatccgtaca tcctgctggt 480
129 gtcgtcgaag gtctcgaccg tcaaggacct gcttcccttg ctggagaagg tcattcagtc 540
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133 ggtc 604
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137 <211> LENGTH: 604
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139 <213> ORGANISM: Mycobacterium avium
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146 gggcctgcgc aacgtcgcgg ccggcgccaa cccgctgggt ctcaagcgcg gcatcgagaa 180
148 ggccgtcgag aaggtcaccg agaccctgct caagtcggcc aaggaggtcg agaccaagga 240
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152 cgaggcgatg gacaaggtcg gcaacgaggg cgtcatcacc gtcgaggagt ccaacacctt 360
154 cggcctgcag ctcgagctca ccgagggtat gcggttcgac aagggttaca tctcgggcta 420
156 cttcgtcacc gacgccgagc gtcaggaagc cgtcctcgag gatccgttca tcctgctggt 480
158 cagctccaag gtctcgaccg tcaaggacct gctgccgctg ctggagaagg tcatccaggc 540

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166 <211> LENGTH: 604
167 <212> TYPE: DNA
168 <213> ORGANISM: Mycobacterium bovis
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175 gggcctgcgc aacgtcgcgg ccggcgccaa cccgctcggt ctcaaacgcg gcatcgaaaa      180
177 ggccgtggag aaggtcaccg agaccctgct caagggcgcc aaggaggtcg agaccaagga      240
179 gcagattgcg gccaccgcag cgatttcggc gggtgaccag tccatcggtg acctgatcgc      300
181 cgaggcgatg gacaaggtgg gcaacgaggg cgtcatcacc gtcgaggagt ccaacacctt      360
183 tgggctgcag ctcgagctca ccgagggtat gcggttcgac aagggctaca tctcggggta      420
185 cttcgtgacc gacccggagc gtcaggaggc ggtcctggag gacccctaca tcctgctggt      480
187 cagctccaag gtgtccactg tcaaggatct gctgccgctg ctcgagaagg tcatcggagc      540
189 cggtaagccg ctgctgatca tcgccgagga cgtcgagggc gaggcgctgt ccaccctggt      600
191 cgtc                                                                  604
194 <210> SEQ ID NO: 7
195 <211> LENGTH: 604
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197 <213> ORGANISM: Mycobaterium bovis BCG
199 <400> SEQUENCE: 7
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204 gggcctgcgc aacgtcgcgg ccggcgccaa cccgctcggt ctcaaacgcg gcatcgaaaa      180
206 ggccgtggag aaggtcaccg agaccctgct caagggcgcc aaggaggtcg agaccaagga      240
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210 cgaggcgatg gacaaggtgg gcaacgaggg cgtcatcacc gtcgaggagt ccaacacctt      360
212 tgggctgcag ctcgagctca ccgagggtat gcggttcgac aagggctaca tctcggggta      420
214 cttcgtgacc gacccggagc gtcaggaggc ggtcctggag gacccctaca tcctgctggt      480
216 cagctccaag gtgtccactg tcaaggatct gctgccgctg ctcgagaagg tcatcggagc      540
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224 <211> LENGTH: 604
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233 gggcctgcgc aacgtcgccg ccggcgccaa cccgctcggc ctgaagcgcg gcatcgagaa      180
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239 cgaggccatg gacaaggtcg gcaacgaggg cgtcatcacc gtcgaggagt ccaacacctt      360
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255 <213> ORGANISM: Mycobaterium celatum TypeII
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268 cgaggcgatg gacaaggtcg gcaacgaggg cgtcatcacc gtcgaggagt ccaacacctt 360
270 cggcctgcag ctcgagctca ccgagggtat gcgcttcgac aagggctaca tctcgggtta 420
272 cttcgtcacc gacgccgagc gtcaggaggc ggtgctcgag gagccctaca tcctgctggt 480
274 cagctccaag gtgtcgacgg tcaaggatct gctgccgctg ctggagaagg tcatccaggc 540
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278 cgtc 604
281 <210> SEQ ID NO: 10
282 <211> LENGTH: 604
283 <212> TYPE: DNA
284 <213> ORGANISM: Mycobacterium chelonae
286 <400> SEQUENCE: 10
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291 aggtctgcgt aacgtcgctg ccggcgccaa cccgctcggc ctgaagcgcg gcatcgagaa 180
293 ggccgtggag gccgtcacca gctctctgct ggactccgcc aaggagatcg acaccaagga 240
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Input Set : **A:\280277 Sequence Listings.txt**
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Input Set : **A:\280277 Sequence Listings.txt**
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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date